

#### **REPLACEMENT SHEET 1/4**

Coding sequence for the *cbh1* gene (SEQ ID NO: 4). Small case letters represent the signal sequence, large case letters the catalytic domain, bolded italics the linker region, and large case underlined the cellulose-binding domain.

at gtateggaagttggccgtcatcteggccttcttggccacagctcgtgctCAGTCGGCCTGCACTCTCCAATCGGAACTCACGTCATCGACGCCAACTGGCGCTGGACTCACGCTACGAACAGCACCACGAACTGCTACGATGG CAACACTTGGAGCTCGACCCTATGTCCTGACAACGAGACCTGCGCGAAGAACTGCTGTCTGGA CGGTGCCGCCTACGCGTCCACGTACGGAGTTACCACGAGCGGTAACAGCCTCTCCATTGGCTT CTACCAGGAATTCACCCTGCTTGGCAACGAGTTCTCTTTCGATGTTGATGTTTCGCAGCTGCCG TGCGGCTTGAACGGAGCTCTCTACTTCGTGTCCATGGACGCGGATGGTGGCGTGAGCAAGTAT CCCACCAACACCGCTGGCGCCAAGTACGGCACGGGGTACTGTGACAGCCAGTGTCCCCGCGA TCTGAAGTTCATCAATGGCCAGGCCAACGTTGAGGGCTGGGAGCCGTCATCCAACAACGCGA ACACGGCATTGGAGGACACGGAAGCTGCTCTGAGATGGATATCTGGGAGGCCAACTCC ATCTCCGAGGCTCTTACCCCCCACCCTTGCACGACTGTCGGCCAGGAGATCTGCGAGGGTGAT GGGTGCGGCGGAACTTACTCCGATAACAGATATGGCGGCACTTGCGATCCCGATGGCTGCGA CTGGAACCCATACCGCCTGGGCAACACCAGCTTCTACGGCCCTGGCTCAAGCTTTACCCTCGA TACCACCAAGAAATTGACCGTTGTCACCCAGTTCGAGACGTCGGGTGCCATCAACCGATACTA TGTCCAGAATGGCGTCACTTTCCAGCAGCCCAACGCCGAGCTTGGTAGTTACTCTGGCAACGA GCTCAACGATGATTACTGCACAGCTGAGGAGGCAGAATTCGGCGGATCCTCTTTCTCAGACAA GGGCGGCCTGACTCAGTTCAAGAAGGCTACCTCTGGCGGCATGGTTCTGGTCATGAGTCTGTG GGATGATTACTACGCCAACATGCTGTGGCTGGACTCCACCTACCCGACAAACGAGACCTCCTC CACACCGGTGCCGTGCGCGAAGCTGCTCCACCAGCTCCGGTGTCCCTGCTCAGGTCGAATC TCAGTCTCCCAACGCCAAGGTCACCTTCTCCAACATCAAGTTCGGACCCATTGGCAGCACCGG  $AGCCACTACCACTGGAAGCTCTCCCGGACCT\underline{ACCCAGTCTCACTACGGCCAGTGCGGCGGTATT}$ GGCTACAGCGGCCCACGGTCTGCGCCAGCGGCACAACTTGCCAGGTCCTGAACCCTTACTAC TCTCAGTGCCTGTAAAGCTCC



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### **REPLACEMENT SHEET 2/4**

SDS-PAGE western blot using an anti-CBHI showing the reduction in molecular weight of rCBHI expression clones as a function of the introduction of N>A modifications.

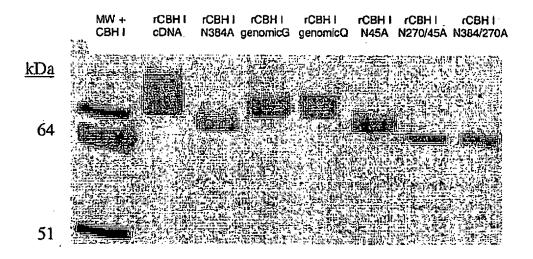


Figure 2



### **REPLACEMENT SHEET 3/4**

## Plasmid map of fungal expression vector pPFE2/CBHI

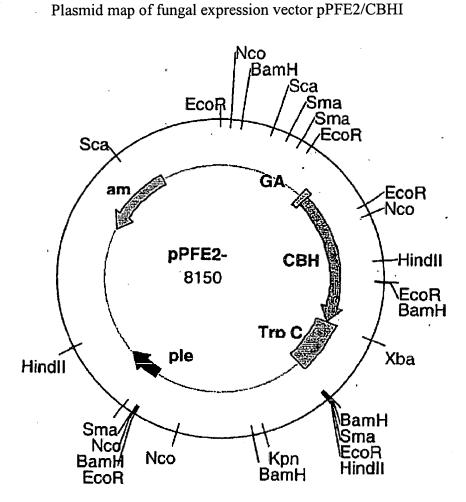


Figure 3



#### **REPLACEMENT SHEET 4/4**

Coding sequence, SEQ ID NO: 19, for the linker region of the *cbh1* gene, SEQ ID NO: 4, showing additional proline residues that effect conformation of the linker region in the protein structure.

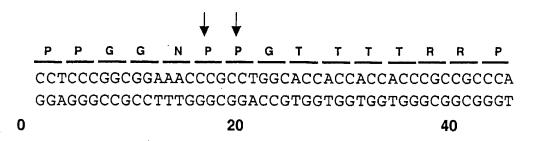


Figure 4